

MAJORITY	SFPKVRAWIEKTL	EEGRRRGYVETL	FGRRRYVPDL	NARVKS	VREAAERMA	FNPVQGT	AADLMK	LAMVKL	
TAQ PRO	768
TFL PRO	767
TTH PRO	770
MAJORITY	FPRLXEMGARM	LQVHDEL	VLEAPKXRAE	XVAALAKE	VM	EGVYPLA	VPLEVE	VGXGEDWLSAKEX	
TAQ PRO	833
TFL PRO	831
TTH PRO	835

FIG. 3C

Genes for Wild-Type and Pol(-)DNAPTaq

Domain
Coding Regions: 5' Nuclease

Polymerase

FIG. 4A

(wt)

FIG. 4B

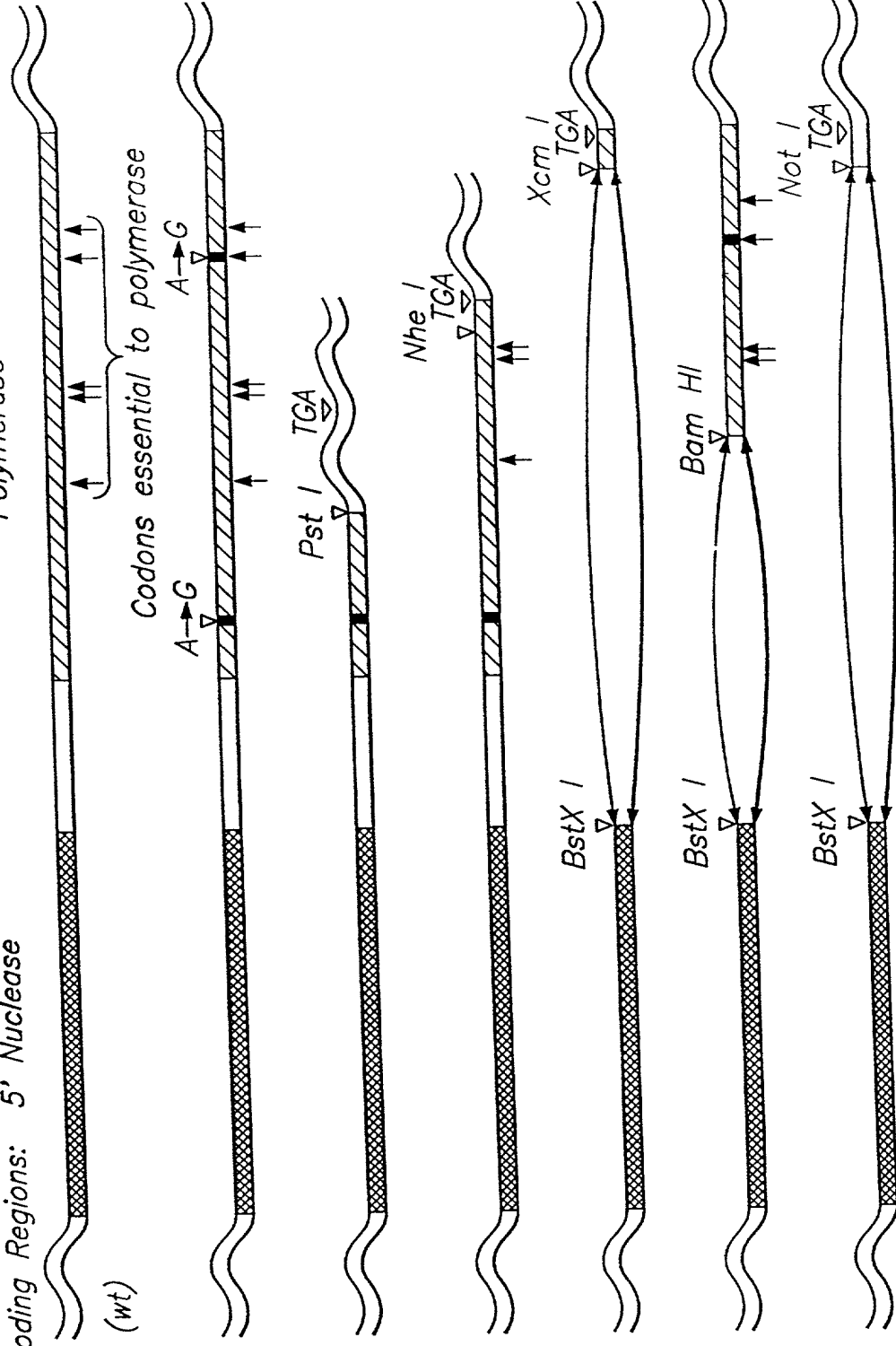
FIG. 4C

FIG. 4D

FIG. 4E

FIG. 4F

FIG. 4G



Genes for Wild-Type and Pol(-)DNAPTfl

